

FIG. 1

1 50
AAAGTCGCAC CTTTCCCCAT AAACCCCTC ACCCCCT TGGACATTGT
51 100
TCCACTTTTC ACTTGATTG TGAAGCACCC AATGCTAGCC CATAGAACAG
101 150
TCCAGTAGTT CAATAGAGAG ACTAGTGAAC ATAGTTTATA ACATTGTCCA
151 200
AGGGGTGGAG GGGGATGCGC GAAATCGATG TGCACGTTTG GTCAAAGATG
201 250
CTCGCGAAAG CTGCACATCA ATTTTCGCACA TGGGCGAAAT TGAATTGCAG
251 300
GTGGGTATAA AAGTTGATGT AGGCCATGTG GCTCGATTTC AACCATATGG
301 350
GTATGCTTCT GAGGATGGGG TGTTACAGTG GACCATATGA GGTAGGTCAT
351 400
TTGAGATGT CACCAAAATG GTCTAAATCT GCGCATTCCA TTTAAGTGAA
401 450
TTTAAGTGAA ATTTAAGTGA ATTTTACTTA AAATTGACCT TTTTCGTTGC
451 500
GCAGATTGG GGTGGTGATG GGTGACGCGG CGAATTTTTT AAAAAAGAGG
500 550
TATATCGCGT GCTATTTGTA TTTTGGTAT CACCGCGTCA CCAATCACCA
551 600
TTGACGGTTT CTTTTTCGAA GTTTTCCGG ATTATTGCAT TTTTATATA
600 650
ATTGTGGGTG GCTGATTCTT GCGAAAGGAC TGTGTGATG TCCGAGTTCC
651 700
CAAATTGGGA GTTTTGGAC ATCACTCCTG ATCTGCCGGC GGCGATCAGG
700 750
ATGACTGACA TTTCGATATA TTTTGGGTAT TCGATAGCTG CCAAATCGGT
751 800
CAGCGTCGAG TATTCGGTT TATTCGAAGG ATTCATGATA TTGCAAAATA
800 850
TCATTGATTT TCATGGGGTT TTGTATTAGT ACCCGCTCAT TGTGGGAAAG
851 900
TCGGGTGGAT TTATCTTACC CGCAAATCTA ATACAAGATT TGCATGATGC
900 950
AGCAATAGAC CAAGGTTAGT ATAGCAGTTG TATTTATACG ACTAGTTATG
951 1000
CAAACCCTTT GTGTTTTTTG TTGCGACTCT TGGCGTGAAC CGGAAGACCG
1000 1050
GACCTCGCTT TCGACTATTC ATCTTTGATG GATATGAGAT CGCAAGGGTA
1051 1100
TCGCTTCGTG CGATATTTAG TGACCATCAG AGCACGCTAC GACTTTTGAT
1100 1150
TATATCCTTG GATTTAATCG GAAGCTCGCA AGCATTGCAT TGATGCAATC

FIG. 2

#1 CACTAGAGTT TCAACAACAT TCGGATTGA CAACATGTCA ACAATTCACA
 #51 ACAGAAATTG ACAACATTGT CACAAATTCT CAAATTGGAC AACATTGGAC
 #101 AAAAATTCAC AACATACATT GGACAACAGT GGACAACGAA CCCAAACCCG
 #151 ACAACATTGT CCAGGGGGAT AGGGGGTGAA AAAGCAGTGC CGGCAAAGTC
 #201 GAAAGATGTC AAGTTGGAAT GCGGCTCAA TTCGTCATTT GTGTAAATCC
 #251 GCAATTTTGC CAATGTGCAA TTTTGCAAAT GTGCAATTTT GCAATGTGC
 #301 AATTTTGCCA ATGTGCAATT TTGCAAATGC GCAATTTTGC AAATCCGCAA
 #351 TTTTGCAAAT GTGCAATTTT GGAAAATCAC CAAATGAAAA TCGTCCAAGT
 #401 CGAATTGGAG GCGTGGTGAC ATGGTCCCAGG GATCCCCTGG TTACAGTGGA
 #451 CAATATCCCA GCAATATTCG CTGTAATTG GAGTTTCGCT GTTTTGCCAA
 #501 ATTTTGAGTC TGAAAAAAA AATTGCAAAT GCGCAAAGGG GGTGAAGGAA
 #551 AAAAAGCAC CCCCGAAGGT AAAATCCCT TTAAGTCCCT TGCGCATTG
 #601 CAAAATTTTC AAAAATTGTT GCAAATGCGC TTTTGTATT TGGCCGGTTC
 #651 ATTGGTGTCA AAAGTTGCCT GGGGTGGTTA CACAATGCAC GGAATTGGTT
 #701 GGAAGTTGTG TGATTGAAAA TTGGTCTGT CACACAATTT TCGGCATTG
 #751 CAAAATTCG CAAATTGGAC AAAAAGGGT CGCGCACAGT CAAATGCGC
 #801 AAATTTCACT TTGAAGTGAG TGCGCATTG TGGGGCAGAA ATGTGGTGAC
 #851 AGCATCGTTT TTTATAATAA ATATTCTATA TTAGTATCT TTATTATAAT
 #901 TTGCTGTAC CAATCACCAT TTTAGAATTT TTATTTTTTT ATGTTTTAGT
 #951 GACCGCGGGA TTTTTGCAA AGTACTATYG TGATGTTGA GTTGTGAA
 #1001 ATGGGCAATT TAGAACATCA TCAGAAATCG CTGAATAGTG ATTTTGAAGT
 #1051 TTGACTGTTT GAAGTGTTTT GGGTATTCGG CAGCTGCCAA ATCGGTCAGC
 #1101 GTCGAATATA ATAGCATTTT TGTGTGTATA TGATATTTAG CGATATCATT
 #1151 GGAATCATGG GGTTTTGTAT TAGTACCCGC TCATTGTGGG AATGTCGGG
 #1201 GGTTCATAT CACCTGCAA TTTAATACAG GATTGTCATG ATGCAGCGAC
 #1251 TGACCGGGGT TGGTATAATA GCTGATTATT CGGCTTATTA TGCAGACCTA
 #1301 TCGTGTAGT AGTTGCGACT CTTGGCGTGA ACCGGAAGAC CGGAAGTGA
 #1351 ATTCGACTAT TTACGTCCGT AACAGGAGA TTTCAAGAAT ATTGCACATT
 #1401 TTGCGTGATA TAAACGTGAT CATCTGAGCA CGCTTCGACT CTTGGATATC
 #1451 TGCTAATCAG CCGTCATCTG AGAGCTCGCA AGCATTGCAA TTGATGCAAT
 #1501

FIG. 3

1	CGTGCCCTTT	TCACGAATTC	ACAGCCCCGC	CCCATGTA	CAATGTTGCC	50
51	CACCCGAAAT	GCCTGCCTGC	CCACCCGAAA	TGCCCGAAAT	GCCCGTTAGA	100
101	AAAAGTATGC	GAAAAGTTCT	TGTCAATTTT	GACAGTGTGT	GAAAAAAGT	150
151	AAAAAGTCCA	CTCAACATTG	CATTATGCAA	TTTGCCACTC	AACATTGTCC	200
201	AGGGGGATAG	GGGGTGAAAA	AGTATCGCAG	TCCAACTGAA	AAGATGCTAA	250
251	GTTGAAATGC	GGCGCAAATT	CATCACTTGA	GTTGCGAAAA	TCCCTAAAGT	300
301	CGAATTGGGC	ACTCGGTGAC	ATGATCGGGA	ATTTCCCTGG	TTACAGTGGT	350
351	CAAATCCCG	CAATTTTGGC	AAAGTTTTTG	AGTTTCGCAC	TTTTTCGAAA	400
401	TTTCGTGTCT	GAAAAAATA	TTTCAACTTT	GCGCAAAGGG	GTCAAAGGGA	450
451	AAAAAAGCAC	CCTCAAAAGG	AAATTTCCCT	TTAATCCCCT	TTGAAAAAAA	500
500	TGCGCAAAGT	TAAATTGCG	AAAATTTGCG	TTTTCTCATA	TGACCGATTA	550
551	GTTGGTGCCA	GATGGTAGTC	GGGATGGTTA	CACGGTGCAC	GGAACGCTT	600
600	GGAAGTTCTG	GAGTTACGAA	TTGGTCCCGT	CACCACAATT	TGCGCATTTT	650
651	TGAAATTGCG	CAAATTGCG	AAAAAAGCAG	CGCGCAAAGT	TAAATTGTGC	700
700	GAAAATTGAC	TTTCAGGTCG	GTGCGCAAAT	TTGGGGTGAA	AAAGTGGTGA	750
751	CAGCATCAGA	ATTATAATA	ATAATCTATA	ATCTAGTTCT	TTTATTATAA	800
800	TTAGCTGTCA	CCAATCACCA	TTTGAGATTT	TTTATTTTTT	TATGTTTTAG	850
851	TGACCGCGGT	ATTTTTTCCA	GAGTACTATC	GTGATGTCTG	AGTTGTCTAA	900
900	AACGGCAATT	TCAGAACATT	ACCAGAAAAC	ACTGAATAGT	GGTTTCTGAG	950
951	TCTGACTGTT	TGAAGTGTTT	TGGGTATTCG	GCAGCTGCCA	ATTCGGTCAG	1000
1000	GGTTGAATAT	ACTAACATTT	CTGTGTGTAT	ATGGTATTTA	GCGATATCAT	1050
1051	TGGAATCATG	GGGTTTTGTA	TTAGTACCCG	CTCATTGTGG	GAAAGTCGGG	1100
1100	TGGTTCAATA	TCACCTGCAA	ATTTAATACA	GGATTGTCAT	GATGCAGCGA	1150
1151	CTGACCGGGG	TTAGTATAAT	AGCTGATTAT	TCGGCTTATT	ATGCAGACCT	1200
1200	ATCGTGTTAG	TAGTTGCGAC	TCTTGCGGTG	AACCGGAAGA	CCGGAAGTTG	1250
1251	ATTTTCGACTA	TTTACGTCCG	TAACACGTCC	GTAAACAGGA	GATTTCAGGA	1300
1300	ATATTGCACA	TTTTGTGTGA	TATAATCGTG	ATCATCTGAG	CACGCTTCGA	1350
1351	CTCTTGAATA	TTTGTAAAC	AACCGATATT	CGGGAGCTCG	CAAGCATTGC	1400
1400	AATTGATGCA	ATC				1450

FIG. 4

Primer	Sequence	Target
300 F	5'-CACTTGTATTGTGAAGCACCC-3'	
300 R	5'-TTG GTG ACA TCT CCA AAT GAC-3'	<i>Perkinsus marinus</i>
500 F	5'-ATGCTAGCCCATAGAACAGT-3'	
500 R	5'-ATGCTAGCCCACATCACAGC-3'	
NTS7	5'-AAGTCGAATTGGAGGCGTGGTGAC-3'	
NTS6	5'-ATTGTGTAACCACCCCAGGC-3'	<i>Perkinsus andrewsi</i>
PM5	5'-ATGCTAGCCC ATAGAACAGT-3'	<i>P. marinus</i> type I
PM7	5'-CAT CTC CAA ATG ACC TAC CT-3'	<i>P. marinus</i> type I
PM6	5'-ATGCTAGCCC ACATCACAGC-3'	<i>P. marinus</i> type II
PM8	5'-CAT CTC CAA ATG ACC TAC CA-3'	<i>P. marinus</i> type II

FIG. 5

FIG. 6

<u>P.sp.</u>		<u>P.o.</u>		<u>P.a.</u>		<u>P.m.</u>			
M	d	a	d	a	d	a	d	a	M

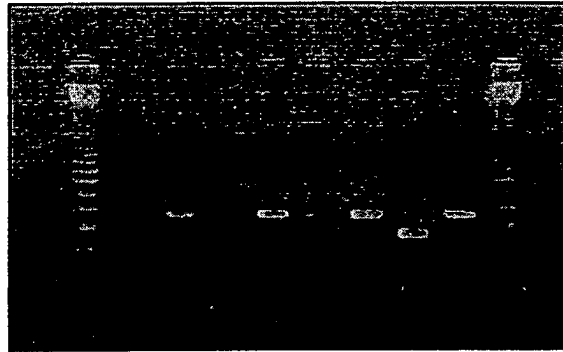


FIG. 7

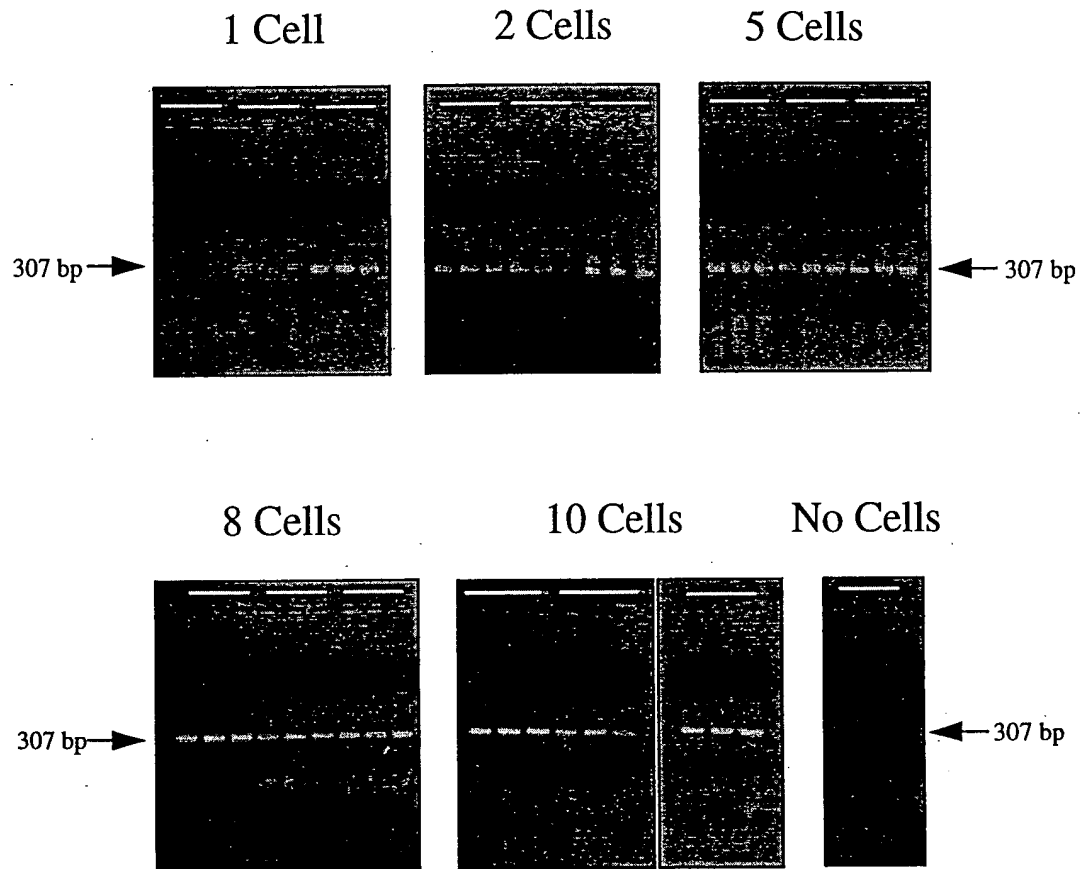


FIG. 8

Samples

1 2 3 4

M a b a b a b a b M

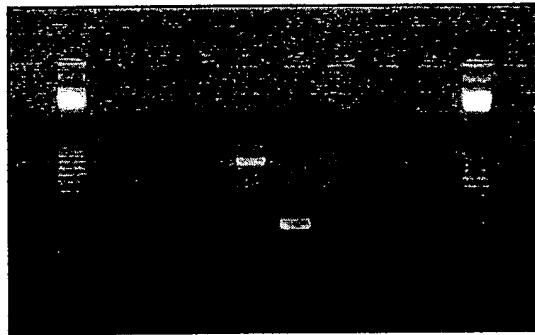


FIG. 9

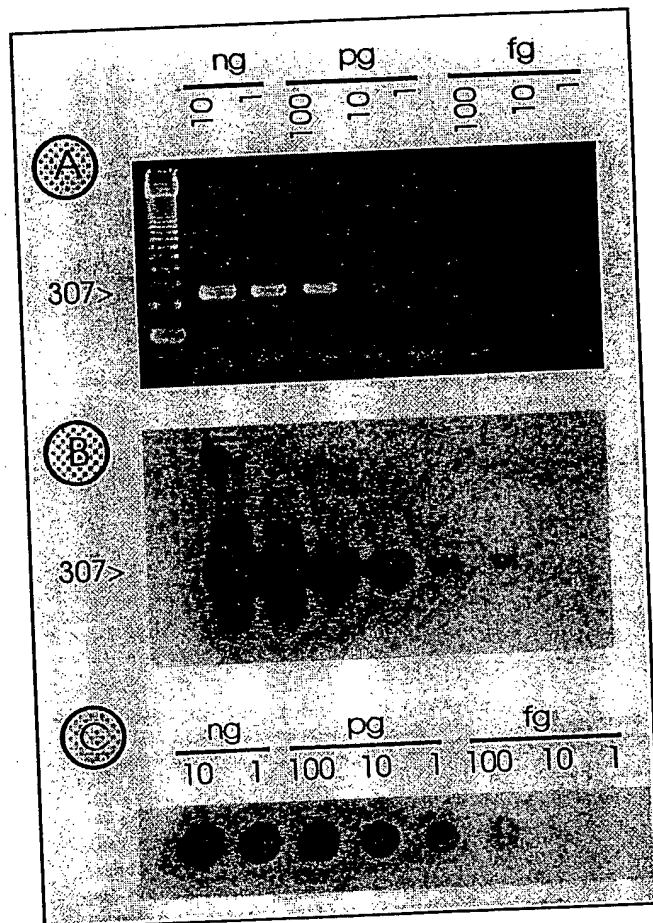


FIG. 10

	1		50
Type-I	CACTTGTATT GTGAAGCACC CAATGCTAGC CCATAGAACA GTCCAGTAGT		
Type-II	CACTTGTATT GTGAAGCACC CAATGCTAGC CCACATCACA GQCCAGTAGT		
	51		100
Type-I	TCAATAGAGA GACTAGTGAA CATAGTTTAT AACATTGTCC AAGGGGTGGA		
Type-II	TCAATAGAGA GACGAGTGAA CATAGTTTAT AACATTGTCC AAGGGGTGGA		
	101		150
Type-I	GGGGGATGCG CGAAATCGAT GTGCACGTTT GGTCAAAGAT GCTCGCGAAA		
Type-II	GGGGGATGCG CGAAATCGAT GTGCACGTTT GGTCAAAGAT GCTCGCGAAA		
	151		200
Type-I	GCTGCACATC AATTTTCGCAC ATGGGCGAAA TTGACTTGCA GGTGGGTATA		
Type-II	GCTGCACATC AATTTTCGCAC ATGGGCGAAA TTGACTTGCA GGTGGGTATA		
	201		250
Type-I	AAAGTTGATG TAGGCCATGT GGCTCGATTT CAACCATATG GGTATGCTTC		
Type-II	AAAGTTGATG TAGGCCATGT GGCTCGATTT CAACCATATG GGTATGCTTC		
	251		300
Type-I	TGAGGATGGG GTGTTACAGT GGACCATATG AGGTAGGTCA TTTGGAGATG		
Type-II	TGAGGATGGG GTGTTACAGT GGACCATATG TGGTAGGTCA TTTGGAGATG		
	301		
Type-I	TCACCAA		
Type-II	TCACCAA		

FIG. 11

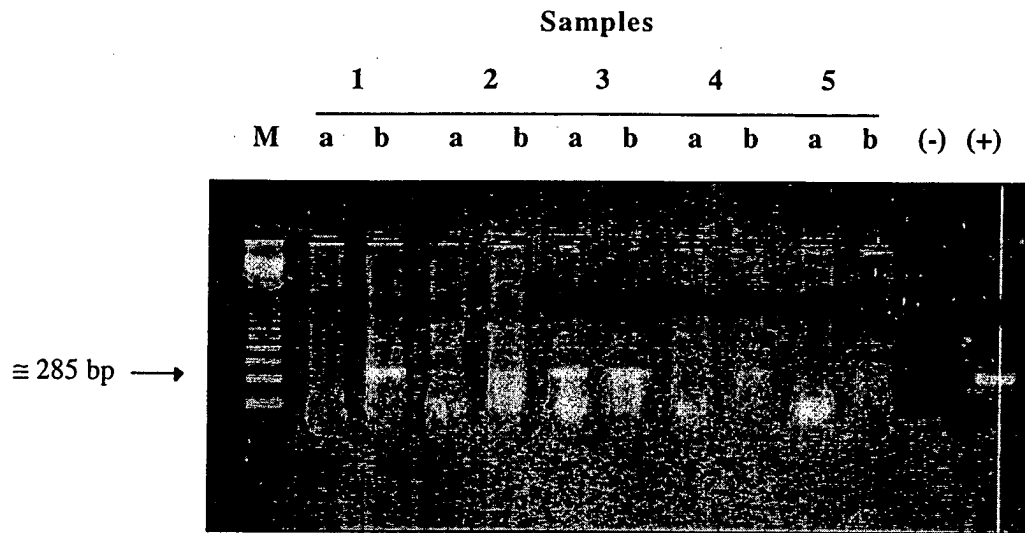


FIG. 12

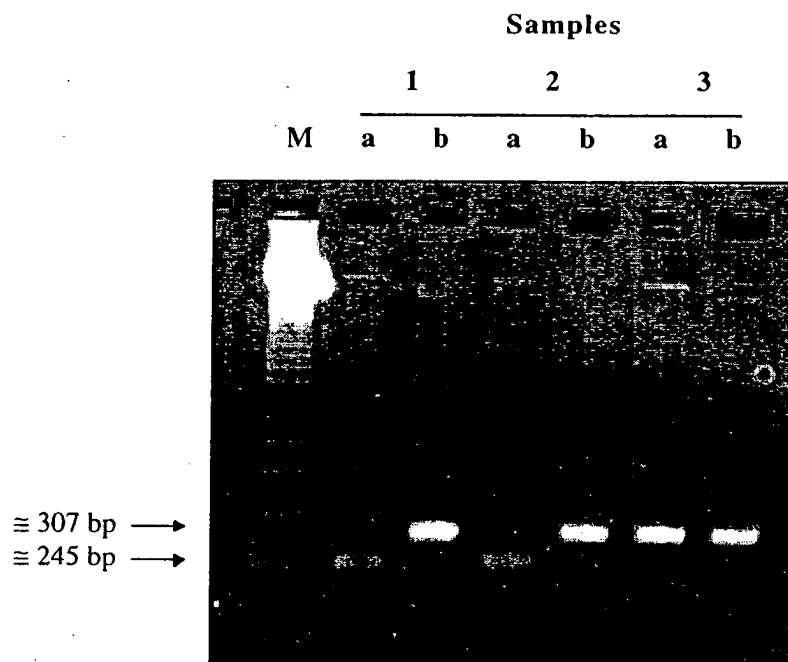


FIG. 13

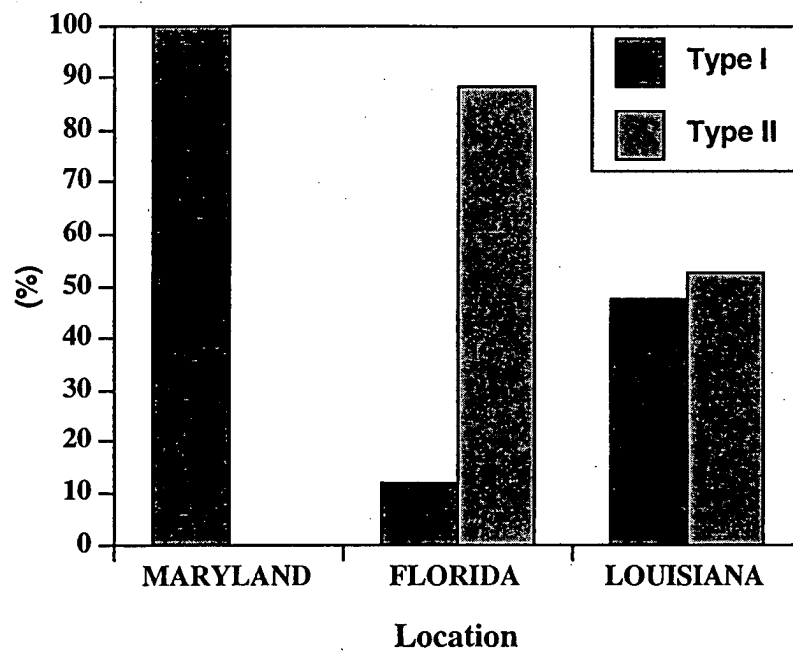


FIG. 14

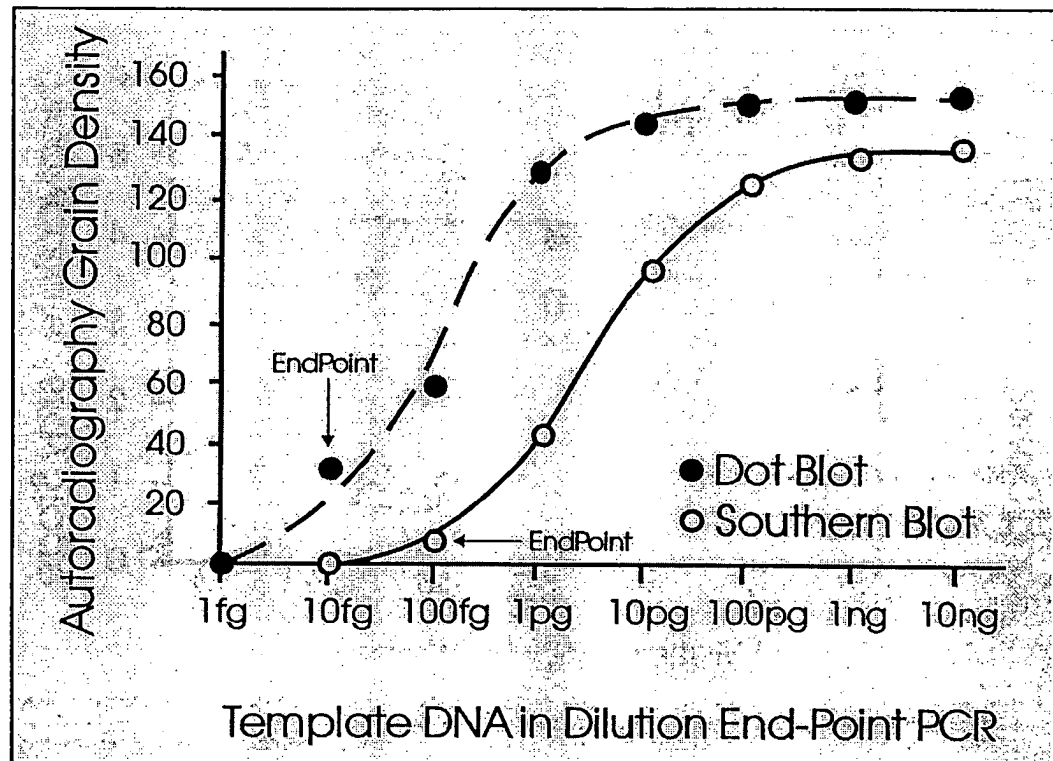
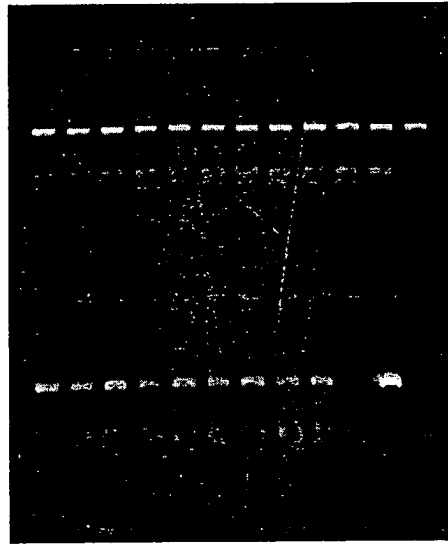


FIG. 15

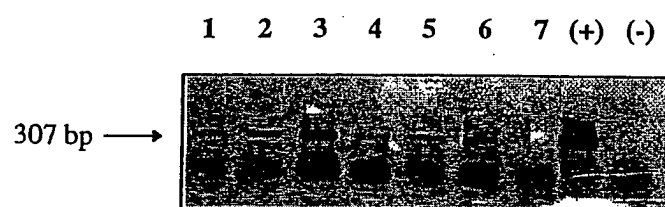
Samples

1 2 3 4 5 6 7 8 9 10 11 12



13 14 15 16 17 18 19 20 1 - + -

FIG. 16



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#1 .TCTTTTAA TCGCACTCAT GGCTTGTGCA TGCAAG CCCCCGAGC
.....
>P. atlanticus.CCCCTGGACA ATGTTATCCC AGCTCAACAA CGAGCAACAG TGCTATGGCA
#51 .....
>P. atlanticus.AGTAGTCCAC TAGAGAGCCA AGTCGACAAT CTCTACAACA TTGTCCAAGG
#101 .....
>P. atlanticus.GGGAAAGGGG GGC GCGCAA GTTGACCTGC AGCAGAGGGA AAAGATGCTG
#151 .....
>P. atlanticus.AGTTTGTCTG CACCCCAACT TTGCGCACTT GGCGAAGTTG ACTTGCAGGC
#201 .....
>P. atlanticus.GAGGGTAAAA GATGCTATGG TTGGTTGCGG ACCAAGTTG CCGTGTGGGT
>PA690F-Text ATGCTATGG TTGGTTGCGG ACC
#251 .....
>P. atlanticus.CATCATTATC GAGGTCTGTG GTGACGATGG ACTAGTTTTT AGGGATTTTT
#301 .....
>P. atlanticus.CGGAGGTGTC ACCACGGACC CCCCAACTTT GCGCACGGGG GGTACTCAAT
#351 .....
>P. atlanticus.TTTAAGTGAA ATTTAAGTAA AATTTACTTA AAATTCACGT TTTTGGGTGC
#401 .....
>P. atlanticus.GCAAAGTTGA GGTGGTGACT GGTGACACGA AAATTTTAAA AAAGAGAGAT
#451 .....
>P. atlanticus.ATTAAAAAAA TATTTATATT TTCTGTGTCA CCGTGTCAAC AGTCACCACA
#501 .....
>P. atlanticus.GGGCGTAATT TTCCGGGAAA TTTTCAGATT TTCCGGAAAA ATTGCATTTT
#551 .....
>P. atlanticus.GGGGTAAATA GTGTCCGTCA GAATTTTGCC AAAGGACTGT CGTGATGTCC
#601 .....
>P. atlanticus.GAGTTCCCAA ATTGAGGGTT TTTGGACATC GCTCTGAAAT CGCTAACGGC
#651 .....
>P. atlanticus.GTTTCAGATT TCCGACTTTT CGACATATTC TGGGTATTTG ATAGCTGCCA
#701 .....
>P. atlanticus.AATCGGTGAG CGTCGAATAT TCCAATATTT CGAAGGATAT ATGATATCGC
#751 .....
>P. atlanticus.GAGATATCAT TGGATTTCAT GGGGTTTTGT ATTAGTACCC GCTCATTGTG
>PER1-Text TAGTACCC GCTCATTGTG
#801 .....
>P. atlanticus.GGAAAGTCGG GTGAATTTAT TCAACCCGCA AATCTAATAC AAGATTGCA
>PER1-Text G
#851 .....
>P. atlanticus.TGATGCAGCG ACTGACCGGG GTGAGTGTAG CAGCTGTTCT ACGGCTTGCT
<PA690R-Text GCTGTTCT ACGGCTTGCT
#901 .....
>P. atlanticus.ACGCAGACCT ATCGTGTTAG TAGTTGCGAC TCTTGCGTG AACCGGAAGA
<PA690R-Text AC
#951 .....
>P. atlanticus.CCGGACCTCG CTTTCGACTA TTCATTCCGA TGAATATGAG ATTGCAAGGG
#1001 .....
>P. atlanticus.TATCGCTTCG TGCGATATTT AGTGATCATC AGAGCAGCT ACGACTTCAG
#1051 .....
>P. atlanticus.TATATCCTCG GATACACAGA AGCTCGCAAG CATTGCATGA TGCAATC
<PER2-Text AGCTCGCAAG CATTGCA
#1101 .....

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FIG. 17

>P. andrewsi-S.ACCTGGTTGA TCCTGCCAGT AGTCATATGC TTGTCTCAAA GATTAAGCCA
#1

>P. andrewsi-S.TGCATGTCTA AGTATAAGCT TTAAACGGCG AAACGCGAA TGGCTCATTA
#51

>P. andrewsi-S.AAACAGTTAT AGTTTATTTG GTGATCGATT ACTATTTGGA TAACCGTAGT
#101

>P. andrewsi-S.AATTCTAGAG CTAATACATG CGTCAAGGCC CGACTTCGGA AGGGCTGCGT
#151

>P. andrewsi-S.TTATTAGATA CAGAACCAAC CTAGCTCCGC CTAGTCCTTG TTGGTGATTG
#201

>P. andrewsi-S.ATAATAACCC GGCGAATCGC ACGGCTTGTC CGGCGATGGA CCATTCAAGT
#251

>P. andrewsi-S.TTCTGACCTA TCAGCTATGG ACGGTAGGGT ATTGGCCTAC CGTGGCGTTG
#301

>P. andrewsi-S.ACGGGTAACG GGAATTAGG GTTCGATTCC GGAGAGGGAG CCTGAGAAAC
#351

>P. andrewsi-S.GACTACCACA TCTAAGGAAG GCAACAGGCG CGCAAATTAC CCAATCCTGA
#401

>P. andrewsi-S.TACAGGGAGG TAGTGACAAG AAATAACAAT ACAGGGCAAT TCTGTCTTGT
#451

>P. andrewsi-S.AATTGGAATG AGTAGATTTT AAATCTCTTT ACGAGTATCA ATTGGAGGGC
#501

>P. andrewsi-S.AAGTCTGGTG CCAGCAGCCG CGGTAATTCC AGCTCCAATA GCGTATATTA
#551

>P. andrewsi-S.AAGTTGTTGC GGTAAAAAG CTCGTAGTTG GATTTCGTCC TTGGGCGACC
>SSU3F-Text AGTTG GATTTCGTCC TTGGGCG
#601

>P. andrewsi-S.GGTCCACCTT TCCTACGGGT TAGGTTGGTA CCAGGTTTGA CCTTGGCTTT
#651

>P. andrewsi-S.TTCTTGGGAT TCGTGCTCAC GCACTTAACT GTGCGCTGAC CGTGTTCCAA
#701

>P. andrewsi-S.GACTTTTACT TTGAGGAAAT TAGAGTGTTT CAAGCAGGCT TATGCCGTGA
#751

>P. andrewsi-S.ATACATTAGC ATGGAATAAT AGGATATGAC TTTGGTCATA TTTGTTGGT
#801

>P. andrewsi-S.TTCTAGGACT GAAGTAATGA TTAATAGGGA CAGTCGGGGG CATTCGTATT
#851

>P. andrewsi-S.TAACTGTCAG AGGTGAAATT CTTGGATTG TTAAAGACGA ACTACTGCGA
#901

FIG.18A

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>P. andrewsi-S.AAGCATTTGC CAAGGATGTT TTCATTGATC AAGAACGAAA GTTAGGGGAT
#951 .....

>P. andrewsi-S.CGAAGACGAT CAGATACCGT CCTAGTCTTA ACCATAAACT ATGCCGACTA
#1001 .....

>P. andrewsi-S.GGGATTGGGA GTCGTTAATT TTAGACGCTC TCAGCACCTC GTGAGAAATC
#1051 .....

>P. andrewsi-S.AAAGTCTTTG GGTTCGGGGG GGAGTATGGT CGCAAGGCTG AAACCTTAAAG
#1101 .....

>P. andrewsi-S.GAATTGACGG AAGGGCACCA CCAGGAGTGG AGCCTGCGGC TTAATTTGAT
#1151 .....

>P. andrewsi-S.TCAACACGGG AAAACTCACC AGGTCCAGAC ATAGGAAGGA TTGACAGATT
>SSU4F-Text ACC AGGTCCAGAC ATAGGAAGG
#1201 .....

>P. andrewsi-S.GATAGCTCTT TCTTGATTCT ATGGGTGGTG GTGCATGGCC GTTCTTAGTT
#1251 .....

>P. andrewsi-S.GGTGGAGTGA TTTGTCTGGT TAATTCGGTT AACGAACGAG ACCTTAACCT
#1301 .....

>P. andrewsi-S.GCTAAATAGT TGCGTGAAAT CTGTATTTC ACCGCTACTT CTTAGAGGGA
#1351 .....

>P. andrewsi-S.CTTTGTGTGT TTAACACAAG GAAGCTTGAG GCAATAACAG GTCTGTGATG
#1401 .....

>P. andrewsi-S.CCCTTAGATG TTCTGGGCTG CACGCGCGCT AACTGACAC GATCAACGAG
#1451 .....

>P. andrewsi-S.TATTCCTTG CCCGGTAGGG TTAGGGTAAT CTTTGAAT CGTGTCGTGC
#1501 .....

>P. andrewsi-S.TAGGGATAGA CGATTGCAAT TATTCGTCTT CAACGAGGAA TTCCTAGTAA
#1551 .....

>P. andrewsi-S.ATGCAAGTCA TCAGCTTGCG TTGATTACGT CCCTGCCCTT TGTACACACC
#1601 .....

>P. andrewsi-S.GCCCGTCGCT CCTACCGATT GAGTGATCCG GTGAGCTGTC CGGACTGCGA
#1651 .....

>P. andrewsi-S.TTAGTTCAGT TTCTGTTCTT TTCGCGGGAA GTTCTGCAA CTTATCACT
#1701 .....

>P. andrewsi-S.TAGAGGAAGG AGAAGTCGTA ACAAGGTTTC CGTAGGTGAA CCTGCAGAAG
#1751 .....

>P. andrewsi-S.GATCATTC

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FIG. 18B

ACACCGATTC ATTCTCTGAG AAACCAGCGG TCTCTGTAAA AGGAGATGGG
#1
ATCTCCGCTT TGTTTAGATC CCCACACCTG ACCGCTTTAA CGGGCCGGGT
#51
AGGTGCATAA CTTCTATGAA CCAATTGTAC TAGTCTAAAG TATCCAATAT
#101
CCTTTTGGAT TTTGGTATTT CAAAACGAAA TTCCAAACTC TCAACGATGG
#151
ATGCCTCGGC TCGAGAATCG ATGAAGGACG CAGCGAAGTG CGATAAGCAC
#201
TGCGATTTGC AGAATTCCGT GAACCAGTAG AAATCTCAAC GCATACTGCA
#251
CAAAGGGGAT TTATCCTCTT TGTACATACA TATCAGTGTC GCTCTTCTTC
#301
CCGATACAAA CATTTTGT TG ATTTACAATC AACATTATGC TTTGTATCCC
#351
GCTTGGATTC CTTTATTGGG ATCCGCTGTG TCGCCTTGCT GACACAGGCG
#401
CATTAAATTTG CAAGGCTATA ATACTACTGT ACTGTAGCCC CTTCGCAAGA
#451
AGGACTGCGC TAGTGAGTAT CTTTGGATGC TCGCGAACTC GACTGTGTTG
#501
TGGTTGATTC CGTGTTCCTC GATCACGCGA TTCATCGCTT CAACGCATTA
#551
TGTCAAATTT GATGAATGCA GAGAGTTGTT TATGAATTAC GCGATCGCTT
#601
TGGTCTCAGA ATCGTTACTA TAGCACGCTT GTCGGTTTGC AACCTGGCAA
#651
TATGTCATCA TT
#701

FIG. 19

<i>Perkinsus</i> species	PCR	Primers to claim							
		Name	Forward Primer (5'-3')	Position ¹	Name	Reverse Primer (5'-3')	Position ¹	Amplicon Size (bp)	Publication
<i>Perkinsus marinus</i>	Species specific	300F	CAC TTG TAT TGT GAA GCA CCC	60-80	300R	TTG GTG ACA TCT CCA AAT GAC	346-366	307	Marsh et al. J. Parasitol. 1995 81(4):577-83. Robledo et al. J. Parasitol. 1999 85(4):650-6.
<i>Perkinsus atlanticus</i>	Species specific	PA690F	ATG CTA TGG TTG GTT GCG GAC C	262-283	PA690R	GTA GCA AGC CGT AGA ACA GC	933-952	691	Robledo et al. J. Parasitol. 2000 86(5):972-8
<i>Perkinsus andrewsi</i> ²	Species specific	NTS7	AAG TCG AAT TGG AGG CGT GGT GAC	447-470	NTS6	ATT GTG TAA CCA CCC CAG CG	717-736	290	Coss et al. J. Euk. Microbiol. 2001 [[In Press]] 48:52- 61
<i>Perkinsus marinus</i>	Generic	PER1	TAG TAC CCG CTC AT(TC) GTG G	827-845	PER2	TGC AAT GCT TGC GAG CT	1123-1139	313	[Coss et al. J. Parasitol. (Submitted)]
<i>Perkinsus atlanticus</i>	Generic	PER1	TAG TAC CCG CTC ATT GTG G	833-851	PER2	TGC AAT GCT TGC GAG CT	1121-1137	305	[Coss et al. J. Parasitol. (Submitted)]
<i>Perkinsus andrewsi</i>	Generic	PER1	TAG TAC CCG CTC ATT GTG G	1121-1239	PER2	TGC AAT GCT TGC GAG CT	1523-1539	319	[Coss et al. J. Parasitol. (Submitted)]

¹Relative to the NTS sequence

²*Perkinsus* sp. (*Macoma balthica*)

FIG. 20

		Primers to claim					
		PCR	Name	Forward Primer (5'-3')	Position	Name	Reverse Primer (5'-3')
<i>Perkinsus</i> species							
<i>Perkinsus andrewsi</i>	Sequencing	SSU3F	SSU4F	AGT TGG ATT TCT GCC TTG CGC G	626-647	1218-1239	ACC AGG TCC AGA CAT AGG AAG G
							Coss et al. J. Euk. Microbiol. 2001 [[In Press]] 48:52- 61

FIG. 21